SEQUENCE LISTING

<110> INCYTE PHARMACEUTICALS, INC.
BANDMAN, Olga
HILLMAN, Jennifer L.
TANG, Y. Tom
LAL, Preeti
CORLEY, Neil C.
GUEGLER, Karl J.
GORGONE, Gina A.
BAUGHN, Mariah R.

<120> HUMAN OXIDOREDUCTASE PROTEINS

<130> PF-0544 PCT

<140> To Be Assigned
<141> Herewith

<150> 60/091,177
<151> 1998-06-30

<160> 14

<170> PERL Program

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Arg Leu Leu Arg Gly Gln Ser Val Gln Gln Val Gly Pro Gln Gly
        35                  40                  45
Leu Leu Tyr Val Gln Gln Arg Glu Leu Ala Val Thr Ser Pro Lys
    50                  55                  60
Asp Gly Ser Ile Ser Ile Leu Gly Ser Asp Asp Ala Thr Thr Cys
65                  70                  75
His Ile Val Val Leu Arg His Thr Gly Asn Gly Ala Thr Cys Leu
                80                  85                  90
Thr His Cys Asp Gly Thr Asp Thr Lys Ala Glu Val Pro Leu Ile
            95                  100                 105
Met Asn Ser Ile Lys Ser Phe Ser Asp His Ala Gln Cys Gly Arg
        110                 115                 120
Leu Glu Val His Leu Val Gly Gly Phe Ser Asp Asp Arg Gln Leu
    125                 130                 135
Ser Gln Lys Leu Thr His Gln Leu Leu Ser Glu Phe Asp Arg Gln
140                 145                 150
Glu Asp Asp Ile His Leu Val Thr Leu Cys Val Thr Glu Leu Asn
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                155                 160                 165
Asp Arg Glu Glu Asn Glu Asn His Phe Pro Val Ile Tyr Gly Ile
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Ala Val Asn Ile Lys Thr Ala Glu Ile Tyr Arg Ala Ser Phe Gln
                185                 190                 195
Asp Arg Gly Pro Glu Glu Gln Leu Arg Ala Ala Arg Thr Leu Ala
                200                 205                 210
Gly Gly Pro Met Ile Ser Ile Tyr Asp Ala Glu Thr Glu Gln Leu
                215                 220                 225
Arg Ile Gly Pro Tyr Ser Trp Thr Pro Phe Pro His Val Asp Phe
                230                 235                 240
Trp Leu His Gln Asp Asp Lys Gln Ile Leu Glu Asn Leu Ser Thr
                245                 250                 255
Ser Pro Leu Ala Glu Pro Pro His Phe Val Glu His Ile Arg Ser
                260                 265                 270
Thr Leu Met Phe Leu Lys Lys His Pro Ser Pro Ala His Thr Leu
                275                 280                 285
Phe Ser Gly Asn Lys Ala Leu Leu Tyr Lys Lys Asn Glu Asp Gly
                290                 295                 300
Leu Trp Glu Lys Ile Ser Ser Pro Gly Ser
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<212> PRT
<213> Homo sapiens

<220>
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<223> Incyte Clone No: 634343

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                20                  25                  30
Tyr Glu His Ala Val Ile Leu Pro Asn Lys Thr Glu Thr Pro Val
                35                  40                  45
Ser Gln Glu Asp Ala Leu Asn Leu Met Asn Glu Asn Ile Asp Ile
                50                  55                  60
Leu Glu Thr Ala Ile Lys Gln Ala Ala Glu Gln Gly Ala Arg Ile
                65                  70                  75
Ile Val Thr Pro Glu Asp Ala Leu Tyr Gly Trp Lys Phe Thr Arg
                80                  85                  90
Glu Thr Val Phe Pro Tyr Leu Glu Asp Ile Pro Asp Pro Gln Val
                95                 100                 105
Asn Trp Ile Pro Cys Gln Asp Pro His Arg Phe Gly His Thr Pro
               110                 115                 120
Val Gln Ala Arg Leu Ser Cys Leu Ala Lys Asp Asn Ser Ile Tyr
               125                 130                 135
Val Leu Ala Asn Leu Gly Asp Lys Lys Pro Cys Asn Ser Arg Asp
               140                 145                 150
Ser Thr Cys Pro Pro Asn Gly Tyr Phe Gln Tyr Asn Thr Asn Val
               155                 160                 165
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Val Tyr Asn Thr Glu Gly Lys Leu Val Ala Arg Tyr His Lys Tyr
                170                 175                 180
His Leu Tyr Ser Glu Pro Gln Phe Asn Val Pro Glu Lys Pro Glu
                185                 190                 195
Leu Val Thr Phe Asn Thr Ala Phe Gly Arg Phe Gly Ile Phe Thr
                200                 205                 210
Cys Phe Asp Ile Phe Phe Tyr Asp Pro Gly Val Thr Leu Val Lys
                215                 220                 225
Asp Phe His Val Asp Thr Ile Leu Phe Pro Thr Ala Trp Met Asn
                230                 235                 240
Val Leu Pro Leu Leu Thr Ala Ile Glu Phe His Ser Ala Trp Ala
                245                 250                 255
Met Gly Met Gly Val Asn Leu Leu Val Ala Asn Thr His His Val
                260                 265                 270
Ser Leu Asn Met Thr Gly Ser Gly Ile Tyr Ala Pro Asn Gly Pro
                275                 280                 285
Lys Val Tyr His Tyr Asp Met Lys Thr Glu Leu Gly Lys Leu Leu
                290                 295                 300
Leu Ser Glu Val Asp Ser His Pro Leu Ser Ser Leu Ala Tyr Pro
                305                 310                 315
Thr Ala Val Asn Trp Asn Ala Tyr Ala Thr Thr Ile Lys Pro Phe
                320                 325                 330
Pro Val Gln Lys Asn Thr Phe Arg Gly Phe Ile Ser Arg Asp Gly
                335                 340                 345
Phe Asn Phe Thr Glu Leu Phe Glu Asn Ala Gly Asn Leu Thr Val
                350                 355                 360
Cys Gln Lys Glu Leu Cys Cys His Leu Ser Tyr Arg Met Leu Gln
                365                 370                 375
Lys Glu Glu Asn Glu Val Tyr Val Leu Gly Ala Phe Thr Gly Leu
                380                 385                 390
His Gly Arg Arg Arg Arg Glu Tyr Trp Gln Val Cys Thr Met Leu
                395                 400                 405
Lys Cys Lys Thr Thr Asn Leu Thr Thr Cys Gly Arg Pro Val Glu
                410                 415                 420
Thr Ala Ser Thr Arg Phe Glu Met Phe Ser Leu Ser Gly Thr Phe
                425                 430                 435
Gly Thr Glu Tyr Val Phe Pro Glu Val Leu Leu Thr Glu Ile His
                440                 445                 450
Leu Ser Pro Gly Lys Phe Glu Val Leu Lys Asp Gly Arg Leu Val
                455                 460                 465
Asn Lys Asn Gly Ser Ser Gly Pro Ile Leu Thr Val Ser Leu Phe
                470                 475                 480
Gly Arg Trp Tyr Thr Lys Asp Ser Leu Tyr Ser Ser Cys Gly Thr
                485                 490                 495
Ser Asn Ser Ala Ile Thr Tyr Leu Leu Ile Phe Ile Leu Leu Met
                500                 505                 510
Ile Ile Ala Leu Gln Asn Ile Val Met Leu
                515                 520
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<210> 3
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<220>
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<223> Incyte Clone No: 1942326

<400> 3

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             20                  25                  30
Asn Phe Val Lys Leu Gln Val Lys Ala Cys Ala Leu Ser Gln Ile
             35                  40                  45
Asn Thr Lys Leu Leu Ala Glu Met Lys Met Lys Lys Asp Leu Phe
             50                  55                  60
Pro Val Gly Arg Glu Ile Ala Gly Ile Val Leu Asp Val Gly Ser
             65                  70                  75
Lys Val Ser Phe Phe Gln Pro Asp Asp Glu Val Val Gly Ile Leu
             80                  85                  90
Pro Leu Asp Ser Glu Asp Pro Gly Leu Cys Glu Val Val Arg Val
             95                 100                 105
His Glu His Tyr Leu Val His Lys Pro Glu Lys Val Thr Trp Thr
            110                 115                 120
Glu Ala Ala Gly Ser Ile Arg Asp Gly Val Arg Ala Tyr Thr Ala
            125                 130                 135
Leu His Tyr Leu Ser His Leu Ser Pro Gly Lys Ser Val Leu Ile
            140                 145                 150
Met Asp Gly Ala Ser Ala Phe Gly Thr Ile Ala Ile Gln Leu Ala
            155                 160                 165
His His Arg Gly Ala Lys Val Ile Ser Thr Ala Cys Ser Leu Glu
            170                 175                 180
Asp Lys Gln Cys Leu Glu Arg Phe Arg Pro Pro Ile Ala Arg Val
            185                 190                 195
Ile Asp Val Ser Asn Gly Lys Val His Val Ala Glu Ser Cys Leu
            200                 205                 210
Glu Glu Thr Gly Gly Leu Gly Val Asp Ile Val Leu Asp Ala Gly
            215                 220                 225
Val Arg Leu Tyr Ser Lys Asp Asp Glu Pro Ala Val Lys Leu Gln
            230                 235                 240
Leu Leu Pro His Lys His Asp Ile Ile Thr Leu Leu Gly Val Gly
            245                 250                 255
Gly His Trp Val Thr Thr Glu Glu Asn Leu Gln Leu Asp Pro Pro
            260                 265                 270
Asp Ser His Cys Leu Phe Leu Lys Gly Ala Thr Leu Ala Phe Leu
            275                 280                 285
Asn Asp Glu Val Trp Asn Leu Ser Asn Val Gln Gln Gly Lys Tyr
            290                 295                 300
Leu Cys Ile Leu Lys Asp Val Met Glu Lys Leu Ser Thr Gly Val
            305                 310                 315
Phe Arg Pro Gln Leu Asp Glu Pro Ile Pro Leu Tyr Glu Ala Lys
            320                 325                 330
Val Ser Met Glu Ala Val Gln Lys Asn Gln Gly Arg Lys Lys Gln
            335                 340                 345
Val Val Gln Phe
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<210> 4
<211> 332
<212> PRT
<213> Homo sapiens

<220>
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<223> Incyte Clone No: 2395269

<400> 4

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Tyr Val Lys Glu Val Ala Lys Pro Ser Pro Gly Glu Gly Glu Val
             20                  25                  30
Leu Leu Lys Val Ala Ala Ser Ala Leu Asn Arg Ala Asp Leu Met
         35                  40                  45
Gln Arg Gln Gly Gln Tyr Asp Pro Pro Pro Gly Ala Ser Asn Ile
     50                  55                  60
Leu Gly Leu Glu Ala Ser Gly His Val Ala Glu Leu Gly Pro Gly
 65                  70                  75
Cys Gln Gly His Trp Lys Ile Gly Asp Thr Ala Met Ala Leu Leu
             80                  85                  90
Pro Gly Gly Gly Gln Ala Gln Tyr Val Thr Val Pro Glu Gly Leu
         95                 100                 105
Leu Met Pro Ile Pro Glu Gly Leu Thr Leu Thr Gln Ala Ala Ala
    110                 115                 120
Ile Pro Glu Ala Trp Leu Thr Ala Phe Gln Leu Leu His Leu Val
    125                 130                 135
Gly Asn Val Gln Ala Gly Asp Tyr Val Leu Ile His Ala Gly Leu
    140                 145                 150
Ser Gly Val Gly Thr Ala Ala Ile Gln Leu Thr Arg Met Ala Gly
    155                 160                 165
Ala Ile Pro Leu Val Thr Ala Gly Ser Gln Lys Lys Leu Gln Met
    170                 175                 180
Ala Glu Lys Leu Gly Ala Ala Ala Gly Phe Asn Tyr Lys Lys Glu
    185                 190                 195
Asp Phe Ser Glu Ala Thr Leu Lys Phe Thr Lys Gly Ala Gly Val
    200                 205                 210
Asn Leu Ile Leu Asp Cys Ile Gly Gly Ser Tyr Trp Glu Lys Asn
    215                 220                 225
Val Asn Cys Leu Ala Leu Asp Gly Arg Trp Val Leu Tyr Gly Leu
    230                 235                 240
Met Gly Gly Gly Asp Ile Asn Gly Pro Leu Phe Ser Lys Leu Leu
    245                 250                 255
Phe Lys Arg Gly Ser Leu Ile Thr Ser Leu Leu Arg Ser Arg Asp
    260                 265                 270
Asn Lys Tyr Lys Gln Met Leu Val Asn Ala Phe Thr Glu Gln Ile
    275                 280                 285
Leu Pro His Phe Ser Thr Glu Gly Pro Gln Arg Leu Leu Pro Val
    290                 295                 300
Leu Asp Arg Ile Tyr Pro Val Thr Glu Ile Gln Glu Ala His Lys
    305                 310                 315
Tyr Met Glu Ala Asn Lys Asn Ile Gly Lys Ile Val Leu Glu Leu
    320                 325                 330
Pro Gln
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<211> 444
<212> PRT
<213> Homo sapiens

<220>
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<223> Incyte Clone No: 008879

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20 25 30
Leu Arg Thr Asp Ser Gly Leu Val Ile Asp Arg Lys Val Tyr Asn
35 40 45
Ile Thr Lys Trp Ser Ile Gln His Pro Gly Gly Gln Arg Val Ile
50 55 60
Gly His Tyr Ala Gly Glu Asp Ala Thr Asp Ala Phe Arg Ala Phe
65 70 75
His Pro Asp Leu Glu Phe Val Gly Lys Phe Leu Lys Pro Leu Leu
80 85 90
Ile Gly Glu Leu Ala Pro Glu Glu Pro Ser Gln Asp His Gly Lys
95 100 105
Asn Ser Lys Ile Thr Glu Asp Phe Arg Ala Leu Arg Lys Thr Ala
110 115 120
Glu Asp Met Asn Leu Phe Lys Thr Asn His Val Phe Phe Leu Leu
125 130 135
Leu Leu Ala His Ile Ile Ala Leu Glu Ser Ile Ala Trp Phe Thr
140 145 150
Val Phe Tyr Phe Gly Asn Gly Trp Ile Pro Thr Leu Ile Thr Ala
155 160 165
Phe Val Leu Ala Thr Ser Gln Ala Gln Ala Gly Trp Leu Gln His
170 175 180
Asp Tyr Gly His Leu Ser Val Tyr Arg Lys Pro Lys Trp Asn His
185 190 195
Leu Val His Lys Phe Val Ile Gly His Leu Lys Gly Ala Ser Ala
200 205 210
Asn Trp Trp Asn His Arg His Phe Gln His His Ala Lys Pro Asn
215 220 225
Ile Phe His Lys Asp Pro Asp Val Asn Met Leu His Val Phe Val
230 235 240
Leu Gly Glu Trp Gln Pro Ile Glu Tyr Gly Lys Lys Lys Leu Lys
245 250 255
Tyr Leu Pro Tyr Asn His Gln His Glu Tyr Phe Phe Leu Ile Gly
260 265 270
Pro Pro Leu Leu Ile Pro Met Tyr Phe Gln Tyr Gln Ile Ile Met
275 280 285
Thr Met Ile Val His Lys Asn Trp Val Asp Leu Ala Trp Ala Val
290 295 300
Ser Tyr Tyr Ile Arg Phe Phe Ile Thr Tyr Ile Pro Phe Tyr Gly
305 310 315
Ile Leu Gly Ala Leu Leu Phe Leu Asn Phe Ile Arg Phe Leu Glu
320 325 330
Ser His Trp Phe Val Trp Val Thr Gln Met Asn His Ile Val Met
335 340 345

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Glu Ile Asp Gln Glu Ala Tyr Arg Asp Trp Phe Ser Ser Gln Leu
                350                 355                 360
Thr Ala Thr Cys Asn Val Glu Gln Ser Phe Phe Asn Asp Trp Phe
                365                 370                 375
Ser Gly His Leu Asn Phe Gln Ile Glu His His Leu Phe Pro Thr
                380                 385                 390
Met Pro Arg His Asn Leu His Lys Ile Ala Pro Leu Val Lys Ser
                395                 400                 405
Leu Cys Ala Lys His Gly Ile Glu Tyr Gln Glu Lys Pro Leu Leu
                410                 415                 420
Arg Ala Leu Leu Asp Ile Ile Arg Ser Leu Lys Lys Ser Gly Lys
                425                 430                 435
Leu Trp Leu Asp Ala Tyr Leu His Lys
                440
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<220>
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                20                  25                  30
Val Leu His Tyr Arg Glu Gly Leu Gly Trp Asp Gly Ser Ala Leu
                35                  40                  45
Glu Phe Asn Trp His Pro Val Leu Met Val Thr Gly Phe Val Phe
                50                  55                  60
Ile Gln Gly Ile Ala Ile Ile Val Tyr Arg Leu Pro Trp Thr Trp
                65                  70                  75
Lys Cys Ser Lys Leu Leu Met Lys Ser Ile His Ala Gly Leu Asn
                80                  85                  90
Ala Val Ala Ala Ile Leu Ala Ile Ile Ser Val Val Ala Val Phe
                95                 100                 105
Glu Asn His Asn Val Asn Asn Ile Ala Asn Met Tyr Ser Leu His
               110                 115                 120
Ser Trp Val Gly Leu Ile Ala Val Ile Cys Tyr Leu Leu Gln Leu
               125                 130                 135
Leu Ser Gly Phe Ser Val Phe Leu Leu Pro Trp Ala Pro Leu Ser
               140                 145                 150
Leu Arg Ala Phe Leu Met Pro Ile His Val Tyr Ser Gly Ile Val
               155                 160                 165
Ile Phe Gly Thr Val Ile Ala Thr Ala Leu Met Gly Leu Thr Glu
               170                 175                 180
Lys Leu Ile Phe Ser Leu Arg Asp Pro Ala Tyr Ser Thr Phe Pro
               185                 190                 195
Pro Glu Gly Val Phe Val Asn Thr Leu Gly Leu Leu Ile Leu Val
               200                 205                 210
Phe Gly Ala Leu Ile Phe Trp Ile Val Thr Arg Pro Gln Trp Lys
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215 220 225
Arg Pro Lys Glu Pro Asn Ser Thr Ile Leu His Pro Asn Gly Gly
230 235 240
Thr Glu Gln Gly Ala Arg Gly Ser Met Pro Ala Tyr Ser Gly Asn
245 250 255
Asn Met Asp Lys Ser Asp Ser Glu Leu Asn Asn Glu Val Ala Ala
260 265 270
Arg Lys Arg Asn Leu Ala Leu Asp Glu Ala Gly Gln Arg Ser Thr
275 280 285
Met

<210> 7
<211> 1126
<212> DNA
<213> Homo sapiens

<220>
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<223> Incyte Clone No: 321510

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tgttcaacaa gtgggacccc agggccttct gtatgttcag caaagagagc ttgcagtgac 180
ctccccaaag gatggctcca tctccattct gggttctgat gatgccacta cttgtcacat 240
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caccaaagct gaggtcccct tgatcatgaa ctccataaaa tccttttctg accacgctca 360
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aaaactcact catcaacttc ttagtgaatt tgacaggcaa gaagatgaca ttcacttagt 480
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tgaagatggc ttgtgggaaa agatctcttc tccaggaagt taaaaaacat gaattaccaa 960
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<212> DNA
<213> Homo sapiens

<220>
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cttacctgac tacttaaaag caaaagagtt aattaagtat tactaattgg tgatactaga 180
tcaatgaaga gaaagcagct gagatccaga ggagtggaag gtccccccttg actaaagcta 240
aatcactaaa ccttggccat ggtcacttcc tcttttccaa tctctgtggc agtttttgcc 300
ctaataaccc tgcaggttgg tactcaggac agttttatag ctgcagtgta tgaacatgct 360
gtcattttgc caaataaaac agaaacacca gtttctcagg aggatgcctt gaatctcatg 420
aacgagaata tagacattct ggagacagcg atcaagcagg cagctgagca gggtgctcga 480
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tatgtcttgg caaatttggg ggacaaaaag ccatgtaatt cccgtgactc cacatgtcct 720
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gctgaatgtg tttatcggct tcccaagttt actaagaaac tttgaagggc tatttcagta 1920
gtatagacca gtgagtccta aatatttttt ctcatcaata attatttttt aagtattatg 1980
ataatgttgt ccatttttttt ggctactctg aaatgttgca gtgtggaaca atggaaagag 2040
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gactttgtgt gtatggggga cttgtatgta tgggagtgag gagtttcagg gccattgcaa 2160
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<210> 9
<211> 1471
<212> DNA
<213> Homo sapiens

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<223> Incyte Clone No: 1942326

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aggaagatct tcctgttaca gaggataact ttgtgaaact tcaagttaaa gcttgtgctc 180
tgagccagat aaatacaaag cttctggcag aaatgaagat gaaaaaggat ttatttcctg 240
ttgggagaga aattgctgga attgtattag atgttggaag caaggtatca ttctttcaac 300
cagatgatga agtagttgga attttgcccc tggactctga agaccctgga ctttgtgaag 360
ttgttagagt acatgagcat tacttggttc ataaaccaga aaaggtcaca tggacggaag 420
cagcaggaag cattcgggat ggagtgcgtg cctatacagc tctgcattat ctttctcatc 480
tctctcctgg aaaatcagtg ctgataatgg atggagcaag tgcatttggt acaatagcta 540

ttcagttagc acatcataga ggagccaaag tgatttcaac agcatgcagc cttgaagata 600
agcagtgcct tgaaagattc agacctccca tagcccgagt gattgatgta tctaatggga 660
aagttcatgt tgctgaaagc tgtttggaag aaacaggtgg cctgggagta gatattgtcc 720
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aaaacaagtt gcatttttaa gcacgtttct ctgctaagac aagatgctca gttgacacat 1260
ttgaaaagtg tttgaaaaat tcttgtgcaa atgatcaaga taattctata attaacatct 1320
taagggaatt tttctaaaaa ccttttcatt gtttctatat attttgcccc tgctataaaa 1380
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gttgcaggcc tttgctaagc taaaaaaaaa a 1471

<210> 10
<211> 1215
<212> DNA
<213> Homo sapiens

<220>
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<223> Incyte Clone No: 2395269

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<213> Homo sapiens

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<212> DNA
<213> Homo sapiens

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<221> misc_feature
<223> GenBank ID No: g1040729

<400> 13

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Asn Asp Leu Trp Ile Ser Ile Leu Gly Lys Val Tyr Asn Val Thr
             35                  40                  45
Glu Trp Ala Lys Glu His Pro Gly Gly Asp Ala Pro Leu Ile Asn
             50                  55                  60
Leu Ala Gly Gln Asp Val Thr Asp Ala Phe Ile Ala Phe His Pro
             65                  70                  75
Gly Thr Ala Trp Lys His Leu Asp Lys Leu Phe Thr Gly Tyr His
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Leu Lys Asp Tyr Gln Val Ser Asp Ile Ser Arg Asp Tyr Arg Lys
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Leu Ala Ser Glu Phe Ala Lys Ala Gly Met Phe Glu Lys Lys Gly
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His Gly Val Ile Tyr Ser Leu Cys Phe Val Ser Leu Leu Leu Ser
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                155                 160                 165
Tyr Leu Gly His Asp Ala Gly His Tyr Gln Met Met Ala Thr Arg
                170                 175                 180
Gly Trp Asn Lys Phe Ala Gly Ile Phe Ile Gly Asn Cys Ile Thr
                185                 190                 195
Gly Ile Ser Ile Ala Trp Trp Lys Trp Thr His Asn Ala His His
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Ile Ala Cys Asn Ser Leu Asp Tyr Asp Pro Asp Leu Gln His Leu
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Pro Met Leu Ala Val Ser Ser Lys Leu Phe Asn Ser Ile Thr Ser
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Val Phe Tyr Gly Arg Gln Leu Thr Phe Asp Pro Leu Ala Arg Phe
                245                 250                 255
Phe Val Ser Tyr Gln His Tyr Leu Tyr Tyr Pro Ile Met Cys Val
                260                 265                 270
Ala Arg Val Asn Leu Tyr Leu Gln Thr Ile Leu Leu Leu Ile Ser
                275                 280                 285
Lys Arg Lys Ile Pro Asp Arg Gly Leu Asn Ile Leu Gly Thr Leu
                290                 295                 300
Ile Phe Trp Thr Trp Phe Pro Leu Leu Val Ser Arg Leu Pro Asn
                305                 310                 315
Trp Pro Glu Arg Val Ala Phe Val Leu Val Ser Phe Cys Val Thr
                320                 325                 330
Gly Ile Gln His Ile Gln Phe Thr Leu Asn His Phe Ser Gly Asp
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Val Tyr Val Gly Pro Pro Lys Gly Asp Asn Trp Phe Glu Lys Gln
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Thr Arg Gly Thr Ile Asp Ile Ala Cys Ser Ser Trp Met Asp Trp
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Arg Leu Pro Arg Cys His Leu Arg Ser Ile Ser Pro Ile Cys Arg
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Glu Leu Cys Lys Lys Tyr Asn Leu Pro Tyr Val Ser Leu Ser Phe
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Tyr Asp Ala Asn Val Thr Thr Leu Lys Thr Leu Arg Thr Ala Ala
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<400> 14

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Ser Asp Leu Gln Phe Asn Ala His Pro Leu Cys Met Val Ile Gly
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Phe Asp Tyr His Arg Lys Lys Gly Tyr Ala Asp Leu Tyr Ser Leu
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Thr Ile Phe Leu Leu Pro Val Gly Thr Ala Leu Leu Gly Leu Lys
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Glu Ala Leu Leu Phe Asn Leu Gly Gly Lys Tyr Ser Ala Phe Glu
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Pro Glu Gly Val Leu Ala Asn Val Leu Gly Leu Leu Leu Ala Cys
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